SEQUENCE LISTING

- <110> OHARA, Osamu NAGASE, Takahiro NOMURA, Nobuo HINUMA, Shuji FUJII, Ryo KITAHARA, Osamu MOGI, Shinichi
- <120> Novel G Protein Coupled Receptor Protein and Its DNA
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660 665 670

Leu Ala Ala Lys Glu Asn Val Val Leu Glu Val Thr Val Leu Asn Thr 680 Glu Gly Gln Val Gln Glu Leu Val Phe Pro Gln Glu Glu Tyr Pro Arg 695 Lys Asn Ser Ile Gln Leu Ser Ala Lys Thr Ile Lys Gln Asn Ser Arg 710 715 Asn Gly Val Val Lys Val Val Phe Ile Leu Tyr Asn Asn Leu Gly Leu 730 Phe Leu Ser Thr Glu Asn Ala Thr Val Lys Leu Ala Gly Glu Ala Gly Pro Gly Gly Pro Gly Gly Ala Ser Leu Val Val Asn Ser Gln Val Ile 760 Ala Ala Ser Ile Asn Lys Glu Ser Ser Arg Val Phe Leu Met Asp Pro 775 Val Ile Phe Thr Val Ala His Leu Glu Asp Lys Asn His Phe Asn Ala 785 790 Asn Cys Ser Phe Trp Asn Tyr Ser Glu Arg Ser Met Leu Gly Tyr Trp 810 Ser Thr Gln Gly Cys Arg Leu Val Glu Ser Asn Lys Thr His Thr Thr 825 Cys Ala Cys Ser His Leu Thr Asn Phe Ala Val Leu Met Ala His Arg Glu Ile Tyr Gln Gly Arg Ile Asn Glu Leu Leu Leu Ser Val Ile Thr Trp Val Gly Ile Val Ile Ser Leu Val Cys Leu Ala Ile Cys Ile Ser 865 870 875 Thr Phe Cys Phe Leu Arg Gly Leu Gln Thr Asp Arg Asn Thr Ile His 890 885 Lys Asn Leu Cys Ile Asn Leu Phe Leu Ala Glu Leu Leu Phe Leu Val 905 Gly Ile Asp Lys Thr Gln Tyr Glu Ile Ala Cys Pro Ile Phe Ala Gly 920 925 Leu Leu His Tyr Phe Phe Leu Ala Ala Phe Ser Trp Leu Cys Leu Glu 930 935 Gly Val His Leu Tyr Leu Leu Leu Val Glu Val Phe Glu Ser Glu Tyr 950 955 Ser Arg Thr Lys Tyr Tyr Leu Gly Gly Tyr Cys Phe Pro Ala Leu

- Val Val Gly Ile Ala Ala Ile Asp Tyr Arg Ser Tyr Gly Thr Glu 980 985 990
- Lys Ala Cys Trp Leu Arg Val Asp Asn Tyr Phe Ile Trp Ser Phe Ile 995 1000 1005
- Gly Pro Val Ser Phe Val Ile Val Val Asn Leu Val Phe Leu Met 1010 1015 1020
- Val Thr Leu His Lys Met Ile Arg Ser Ser Ser Val Leu Lys Pro 1025 1030 1035
- Asp Ser Ser Arg Leu Asp Asn Ile Lys Ser Trp Ala Leu Gly Ala 1040 1045 1050
- Ile Ala Leu Leu Phe Leu Leu Gly Leu Thr Trp Ala Phe Gly Leu 1055 1060 1065
- Leu Phe Ile Asn Lys Glu Ser Val Val Met Ala Tyr Leu Phe Thr 1070 1075 1080
- Thr Phe Asn Ala Phe Gln Gly Val Phe Ile Phe Val Phe His Cys 1085 1090 1095
- Ala Leu Gln Lys Lys Val His Lys Glu Tyr Ser Lys Cys Leu Arg 1100 1105 1110
- His Ser Tyr Cys Cys Ile Arg Ser Pro Pro Gly Gly Thr His Gly 1115 1120 1125
- Ser Leu Lys Thr Ser Ala Met Arg Ser Asn Thr Arg Tyr Tyr Thr 1130 1135 1140
- Gly Thr Gln Ser Arg Ile Arg Arg Met Trp Asn Asp Thr Val Arg 1145 1150 1155
- Lys Gln Thr Glu Ser Ser Phe Met Ala Gly Asp Ile Asn Ser Thr 1160 1165 1170
- Pro Thr Leu Asn Arg Gly Thr Met Gly Asn His Leu Leu Thr Asn 1175 1180 1185
- Pro Val Leu Gln Pro Arg Gly Gly Thr Ser Pro Tyr Asn Thr Leu 1190 1195 1200
- Ile Ala Glu Ser Val Gly Phe Asn Pro Ser Ser Pro Pro Val Phe 1205 1210 1215
- Asn Ser Pro Gly Ser Tyr Arg Glu Pro Lys His Pro Leu Gly Gly 1220 1225 1230
- Arg Glu Ala Cys Gly Met Asp Thr Leu Pro Leu Asn Gly Asn Phe 1235 1240 1245
- Asn Asn Ser Tyr Ser Leu Arg Ser Gly Asp Phe Pro Pro Gly Asp

	1250					1255					1260			
Gly	Gly 1265	Pro	Glu	Pro	Pro	Arg 1270	_	Arg	Asn	Leu	Ala 1275	Asp	Ala	Ala
Ala	Phe 1280	Glu	Lys	Met	Ile	Ile 1285		Glu	Leu	Val	His 1290	Asn	Asn	Leu
Arg	Gly 1295	Ser	Ser	Ser	Ala	Ala 1300	Lys	Gly	Pro	Pro	Pro 1305	Pro	Glu	Pro
Pro	Val 1310	Pro	Pro	Val	Pro	Gly 1315	Gly	Gly	Gly	Glu	Glu 1320	Glu	Ala	Gly
Gly	Pro 1325	Gly	Gly	Ala	Asp	Arg 1330	Ala	Glu	Ile	Glu	Leu 1335	Leu	Tyr	Lys
Ala	Leu 1340	Glu	Glu	Pro	Leu	Leu 1345	Leu	Pro	Arg	Ala	Gln 1350	Ser	Val	Leu
Tyr	Gln 1355	Ser	Asp	Leu	Asp	Glu 1360	Ser	Glu	Ser	Cys	Thr 1365	Ala	Glu	Asp
Gly	Ala 1370	Thr	Ser	Arg	Pro	Leu 1375	Ser	Ser	Pro	Pro	Gly 1380	Arg	Asp	Ser
Leu	Tyr 1385	Ala	Ser	Gly	Ala	Asn 1390	Leu	Arg	Asp	Ser	Pro 1395	Ser	Tyr	Pro
Asp	Ser 1400	Ser	Pro	Glu	Gly	Pro 1405	Ser	Glu	Ala	Leu	Pro 1410	Pro	Pro	Pro
Pro	Ala 1415	Pro	Pro	Gly	Pro	Pro 1420	Glu	Ile	Tyr	Tyr	Thr 1425	Ser	Arg	Pro
Pro	Ala 1430	Leu	Val	Ala	Arg	Asn 1435	Pro	Leu	Gln	Gly	Tyr 1440	Tyr	Gln	Val
Arg	Arg 1445	Pro	Ser	His	Glu	Gly 1450	Tyr	Leu	Ala	Ala	Pro 1455	Gly	Leu	Glu
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Leu														
<210> 6														

<211> 4422 <212> DNA

<213> Homo sapiens

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180 tgtgaagget accecatega getgeggtge ceeggeageg aegteateat ggtggagaat gccaactacg ggcgcacgga cgacaagatt tgcgatgctg accettteca gatggagaat 240 gtgcagtgct acctgccgga cgccttcaag atcatgtcac agaggtgtaa caaccgcacc 300 cagtgcgtgg tggtcgccgg ctcggatgcc tttcctgacc cctgtcctgg gacctacaag 360 420 tacctggagg tgcagtacga ctgtgtcccc tacaaagtgg agcagaaagt cttcgtgtgc 480 ccagggaccc tgcagaaggt gctggagccc acctcgacac acgagtcaga gcaccagtct ggcgcatggt gcaaggaccc gctgcaggcg ggtgaccgca tctacgtgat gccctggatc 540 600 ccctaccgca cggacacact gactgagtat gcctcgtggg aggactacgt ggccgcccgc 660 cacaccacca cctaccgcct gcccaaccgc gtggatggca caggctttgt ggtctacgat ggtgccgtct tctacaacaa ggagcgcacg cgcaacatcg tcaagtatga cctacggacg 720 780 cgcatcaaga gcggggagac ggtcatcaat accgccaact accatgacac ctcgccctac 840 cgctggggcg gaaagaccga cattgacctg gcggtggacg agaacgggct gtgggtcatc 900 tacgccactg agggcaacaa cgggcggctg gtggtgagcc agctgaaccc ctacacactg cgctttgagg gcacgtggga gacgggttac gacaagcgct cggcatccaa cgccttcatg 960 1020 gtgtgtgggg teetgtaegt eetgegetee gtgtaegtgg atgatgaeag egaggegget 1080 ggcaaccgcg tggactatgc cttcaacacc aatgccaacc gcgaggagcc tgtcagcctc 1140 accttcccca acccctacca gttcatctcc tccgttgact acaaccctcg cgacaaccag 1200 ctgtacgtct ggaacaacta tttcgtggtg cgctacagcc tggagttcgg gccgcccgac 1260 cccagtgctg gcccagccac ttccccaccc ctcagcacga ccaccacagc caggcccacg 1320 cccctcacca gcacagcctc gcccgcagcc accaccccgc tccgccgggc acccctcacc acgcacccag tgggtgccat caaccagctg ggacctgatc tgcctccagc cacagcccca 1380 1440 gtccccagca cccggcggcc cccagccccg aatctacacg tgtcccctga gctcttctgc gageceegag aggtaeggeg ggteeagtgg eeggeeacee ageagggeat getggtggag 1500 1560 aggecetgee ceaaggggae tegaggaatt gesteettee agtgtetass agsettgggg 1620 ctctggaacc cccggggccc tgacctcagc aactgcacct ccccctgggt caaccaggtg 1680 gcccagaaga tcaagagtgg ggagaacgcg gccaacatcg ccagcgagct ggcccgacac accoggggct ccatctacgc gggggacgtc tectectetg tgaagetgat ggagcagetg 1740 1800 ctggacatcc tggatgccca gctgcaggcc ctgcggccca tcgagcgcga gtcagccggc 1860 aagaactaca acaagatgca caagcgagag agaacttgta aggattatat caaggccgtg

1920 gtggagacag tggacaatct gctccggcca gaagctctgg agtcctggaa ggacatgaat gccacggagc aggtgcacac ggccaccatg ctcctcgacg tcctggagga gggcgccttc 1980 2040 ctgctggccg acaatgtcag ggagcctgcc cgcttcctgg ctgccaagga gaacgtggtc 2100 ctggaggtca cagtcctgaa cacagagggc caggtgcagg agctggtgtt cccccaggag 2160 gagtaccega gaaagaacte catecagetg tetgecaaaa ecateaagea gaacageege 2220 aatggggtgg tcaaagttgt cttcatcctc tacaacaacc tgggcctctt cctgtccacg gagaatgcca cagtgaaget ggccggcgaa gcaggcccgg gtggccctgg gggcgcctct 2280 2340 ctagtggtga actcacaggt catcgcagca tccatcaaca aggagtccag ccgcgtcttc 2400 ctcatggacc ctgtcatctt caccgtggcc cacctggagg acaagaacca cttcaatgct 2460 aactgctcct tctggaacta ctcggagcgt tccatgctgg gctattggtc gacccaaggc 2520 tgccgcctgg tggagtccaa caagacccat accacgtgtg cctgcagcca cctcaccaac 2580 ttegetgtge teatggetea eegtgagate taceagggee geateaaega getgetgetg 2640 teggteatea cetgggtggg cattgtgate tecetggtet gettggeeat etgeatetee 2700 accttctgct tcctgcgggg gctgcagacc gaccgcaaca ccatccacaa gaacctgtgc 2760 atcaacctct teetggetga getgetette etggteggga tegacaagae teagtatgag attgcctgcc ccatcttcgc cggcctgctg cactatttct tcctggctgc cttctcctgg 2820 2880 ctgtgcctgg agggcgtgca cctctacctg ctactagtgg aggtgtttga gagcgagtat tecegeacea agtactaeta eetgggtgge taetgettee eggeeetggt ggtgggeate 2940 3000 geggetgeea ttgactaceg cagetacgge accgagaagg cetgetgget cegagtggac 3060 aattacttca tetggagttt categggeea gteteetteg ttategtggt caacetggtg 3120 tteeteatgg tgaccetgea caagatgate egaageteat etgtgeteaa geeegaetee 3180 agcogcotgg acaacattaa atcotgggcg otgggggcca togogctgct gttootgctg 3240 ggcctcacct gggctttcgg cctcctcttc atcaacaagg agtcggtggt catggcctat 3300 ctcttcacca ccttcaacgc cttccagggg gtcttcatct tcgtctttca ctgcgcctta cagaagaagg tgcacaagga gtacagcaag tgcctgcgtc actcctactg ctgcatccgc 3360 3420 tececaceeg ggggeactea eggateecte aagaceteag ecatgegaag caacaceege 3480 tactacacag ggacccagag ccgaattcgg aggatgtgga atgacactgt gaggaaacag 3540 acggagtect cetteatgge gggtgacate aacageacce ceaccetgaa cegaggtace

3600 atggggaacc acctgctgac caaccccgtg ctgcagcccc gtgggggcac cagtccctac aacaccctca tegeogagte agtgggette aatccctect egeoccetgt etteaactee 3660 ccagggagct accgggaacc caagcacccc ttgggaggcc gggaagcctg tggcatggac 3720 accetgeece tgaacggeaa etteaataac agttacteet tgegaagtgg ggattteeet 3780 cccggggatg ggggccctga gccgcccga ggccggaacc tagccgatgc ggcggccttt 3840 gagaagatga tcatctcaga gctggtgcac aacaacctgc gggggagcag cagcgcgcc 3900 aagggeeete eacegeetga geeeeetgtg eeacetgtge cagggggegg gggegaggaa 3960 gaggcgggcg ggcccggggg tgctgaccgg gccgagattg aacttctcta taaggccctg 4020 gaggagcete tgetgetgee eegggeeeag teggtgetgt accagagega tetggaegag 4080 4140 teggagaget geaeggeega ggaeggegee accageegge eceteteete eeeteetgge 4200 egggaetece tetatgecag eggggecaae etgegggaet eaccetecta eceggaeage agecetgagg ggeceagtga ggecetgece ceaececete eegeaeeeee eggeeeeeee 4260 gaaatctact acacetegeg ceegecagee etggtggeee ggaateeeet geagggetae 4320 taccaggtgc ggcgtcctag ccacgagggc tacctggcag ccccaggcct tgaggggcca 4380 4422 gggcccgatg gggacgggca gatgcagctg gtcaccagtc tc

<210> 7 <211> 3000

<212> DNA

<213> Homo sapiens

<220>

ᅰ "

<221> misc_feature

<222> (1)..(3000)

<223> Sequence depicted in Figs 1 and 2 inclusive, containing protein e ncoding sequence of seq id two

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gaccagctgg taggcctcct agatgtacag cttcggaact tgaccccagg tggaaaagat 120
agtgctgccc ggagtttgaa caaggcaatg gtcgagacag ttaacaacct ccttcagcca 180
caagctttga atgcatggag agacctgact acgagtgatc agctgcgtgc ggccaccatg 240
ttgcttcata ctgtggagga aagtgcttt gtgctggctg ataacctttt gaagactgac 300
attgtcaggg agaatacaga caatattaaa ttggaagttg caagactgag cacagaagga 360
aacttagaag acctaaaatt tccagaaaac atgggccatg gaagcactat ccagctgtct 420

gcaaatacct taaagcaaaa tggccgaaat ggagagatca gagtggcctt tgtcctgtat 480 aacaacttgg gtccttattt atccacggag aatgccagta tgaagttggg aacggaagct 540 600 ttgtccacaa atcattctgt tattgtcaat tcccctgtta ttacggcagc aataaacaaa 660 gagttcagta acaaggttta tttggctgat cctgtggtat ttactgttaa acatatcaag 720 cagtcagagg aaaatttcaa ccctaactgt tcattttgga gctactccaa gcgtacaatg 780 acaggttatt ggtcaacaca aggctgtcgg ctcctgacaa caaataagac acatactaca tgctcttgta accacctaac aaattttgca gtactgatgg cacatgtgga agttaagcac 840 agtgatgcgg tccatgacct ccttctggat gtgatcacgt gggttggaat tttgctgtcc 900 960 cttgtttgtc tcctgatttg catcttcaca ttttgctttt tccgcgggct ccagagtgac cgtaacacca tccacaagaa cctctgcatc agtctctttg tagcagagct gctcttcctg 1020 1080 attgggatca accgaactga ccaaccaatt gcctgtgctg ttttcgctgc cctgttttct tettettgge tgeetteace tggatgttee tggagggggt geagetttat atacateatg 1140 1200 ctggtggagg tttttgagag tgaacattca cgtaggaaat acttttatct ggtcggctat gggatgcctg cactcattgt ggctgtgtca gctgcagtag actacaggag ttatggaaca 1260 1320 gataaagtat gttggctccg acttgacacc tacttcattt ggagttttat aggaccagca 1380 actttgataa ttatgcttaa tgtaatcttc cttgggattg ctttatataa aatgtttcat catactgcta tactgaaacc tgaatcaggc tgtcttgata acatcaagtc atgggttata 1440 1500 ggtgcaatag ctcttctctg cctattagga ttgacctggg cctttggact catgtatatt 1560 aatgaaagca cagtcatcat ggcctatctc ttcaccattt tcaattctct acagggaatg 1620 tttatattta ttttccattg tgtcctacag aagaaggtac gaaaagagta tgggaaatgc 1680 ctgcgaacac attgctgtag tggcaaaagt acagagagtt ccattggttc agggaaaaca totggttotc gaactootgg acgotactoc acaggotcac agagocgaat cogtagaatg 1740 1800 tggaatgaca cggttcgaaa gcagtcagag tcttccttta ttactggaga cataaacagt 1860 tcagcgtcac tcaacagaga ggggcttctg aacaatgcca gggatacaag tgtcatggat 1920 actictaccae tgaatggtaa ceatggeaat agttacagea ttgccagegg egaatacetg 1980 agcaactgtg tgcaaatcat agaccgtggc tataaccata acgagaccgc cctagagaaa 2040 aagattetga aggaacteae ttecaactat atceettett acetgaacaa ecatgagege 2100 tccagtgaac agaacaggaa tctgatgaac aagctggtga ataaccttgg cagtggaagg

2160 gaagatgatg ccattgtcct ggatgatgcc acctcgttta accacgagga gagtttgggc 2220 ctggaactca ttcatgagga atctgatgct cctttgctgc ccccaagagt atactccacc gagaaccacc agccacacca ttataccaga aggcggatcc cccaagacca cagtgagagc 2280 2340 tttttccctt tgctaaccaa cgagcacaca gaagatctcc agtcacccca tagagactct 2400 ctctatacca gcatgccgac actggctggt gtggccgcca cagagagtgt taccaccagc acccagaccg aacccccacc ggccaaatgt ggtgatgccg aagatgttta ctacaaaagc 2460 atgccaaacc taggctccag aaaccacgtc catcagctgc atacttacta ccagctaggt 2520 cgcggcagca gtgatggatt tatagttcct ccaaacaaag atgggacccc tcccgaggga 2580 2640 agttcaaaag gaccggctca tttggtcact agtctataga agatgacaca gaaattggaa 2700 ccaacaaaac tgctaacacc ttgttgactg ttctgagttg atataagcag tggtaataat 2760 gtgtgtactc ctaaatcttt atgctgtcct ctaaagacaa acacaaactc tcagactttt 2820 ttttttttta atgggatttt taggtcagcc caggggagaa agataactgc taaaattccc 2880 ctgtacccca tcctttcttg tcctttcccc ttcagatgga gacttcatta tgttaatgaa caagatatga agaaaatggc actcattgtg gccttgttga attatgttgt gtatgtttta 2940 3000 acatctctga tgctgtgtta ctaaaattac aaggacctgc tttttaaaag gccagaacaa

<210> 8 <211> 4343 <212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)..(4343)

<223> Sequence depicted in Figs 7-15 inclusive, containing protein encoding sequence of seq id four

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actgcctcat atctctgcat gatttccact ggaacatgga accctaaggg ccccgatctt 240
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gctgctagtc ttgccaatga actggctaaa cataccaaag ggccagtgtt tgctggggat 360
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